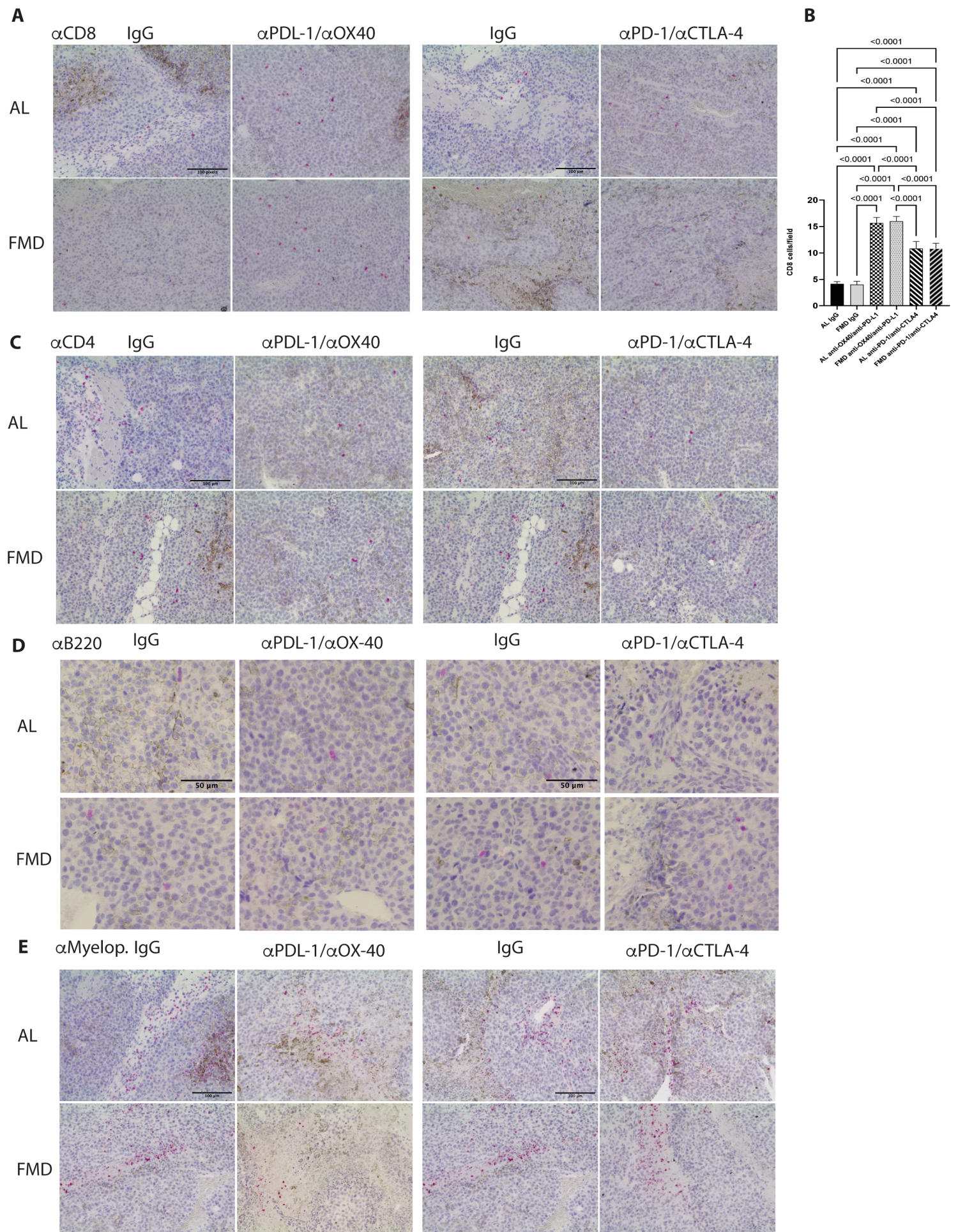
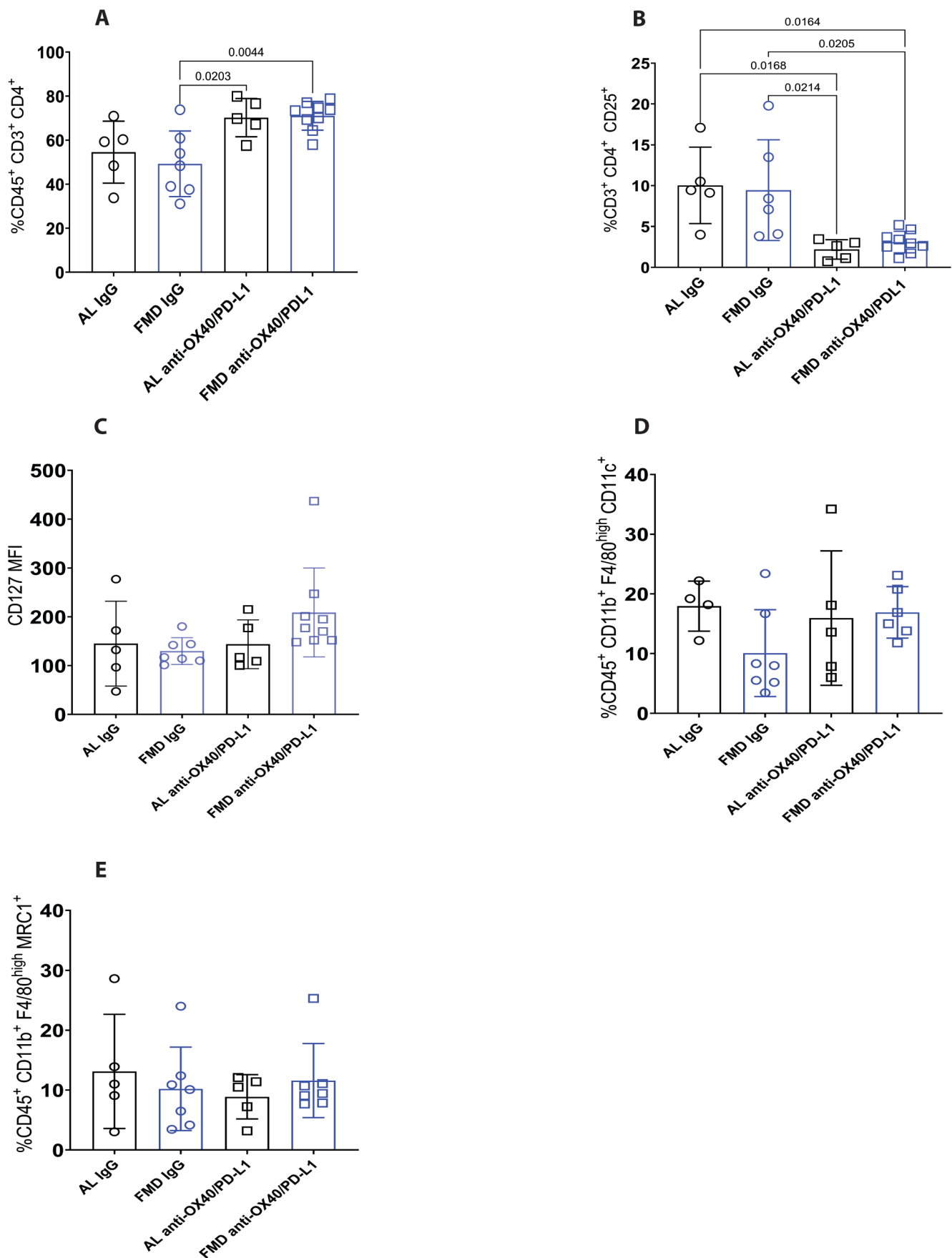


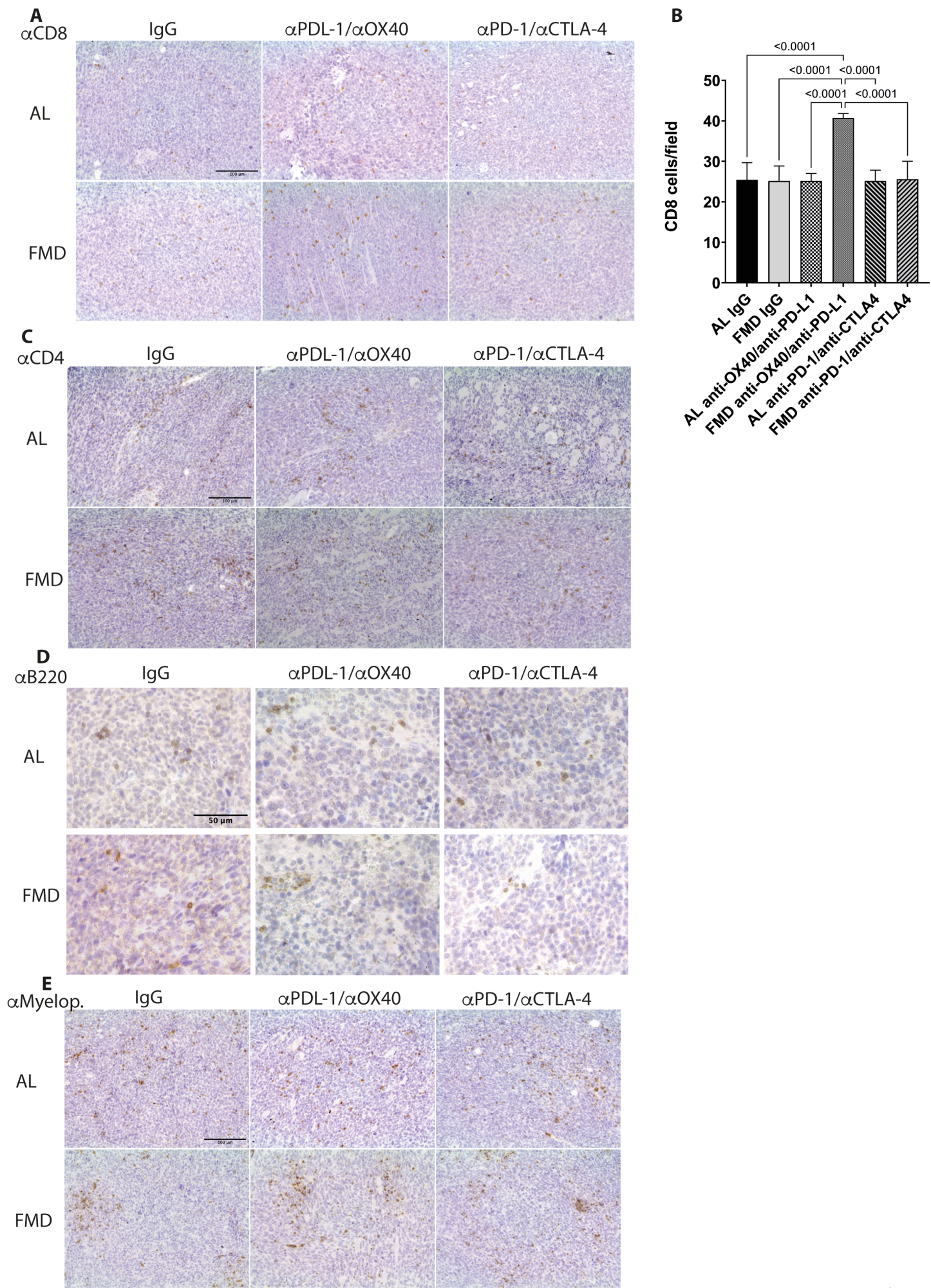
Suppl Figure 1. Analysis of B16 breast tumor infiltrating lymphocytes upon immunotherapy treatment in standard diet and FMD group. **(A)** CD45⁺CD3⁺CD4⁺ T cell (AL NT n=6; FMD NT n=5; AL PD1 CTLA4 n=6; FMD PD1 CTLA4 n=6); **(B)** CD3⁺CD4⁺CD25⁺ Treg cells (AL NT n=6; FMD NT n=5; AL PD1 CTLA4 n=6; FMD PD1 CTLA4 n=6); **(C)** CD45⁺CD11b⁺Ly6C^{high} M-MDSC (AL NT n=6; FMD NT n=4; AL PD1 CTLA4 n=6; FMD PD1 CTLA4 n=5); **(D)** CD45⁺CD11b⁺Ly6C^{low} Ly6G^{high}PMN-MDSC (AL NT n=6; FMD NT n=5; AL PD1 CTLA4 n=6; FMD PD1 CTLA4 n=6); **(E)** CD45⁺ CD11c⁺MHCII⁺ dendritic cell (AL NT n=6; FMD NT n=5; AL PD1 CTLA4 n=6; FMD PD1 CTLA4 n=6); **(F)**CD45⁺ CD11b⁺F4/80^{high} macrophage (AL NT n=6; FMD NT n=5; AL PD1 CTLA4 n=6; FMD PD1 CTLA4 n=6), **(G)** CD45⁺CD11b⁺F480^{high}CD11c⁺ M1 (AL NT n=6; FMD NT n=5; AL PD1 CTLA4 n=6; FMD PD1 CTLA4 n=6); **(H)** CD45⁺CD11b⁺F480^{high}MRC1⁺ M2 macrophage (AL NT n=5; FMD NT n=4; AL PD1 CTLA4 n=6; FMD PD1 CTLA4 n=5). Statistical analysis was performed using one-way analysis of variance (ANOVA). P values were determined by One-way ANOVA with Tukey's post analysis. Differences were considered significant when P≤0.05. All data are represented as mean ± SEM. Source data are provided as a Source Data file.



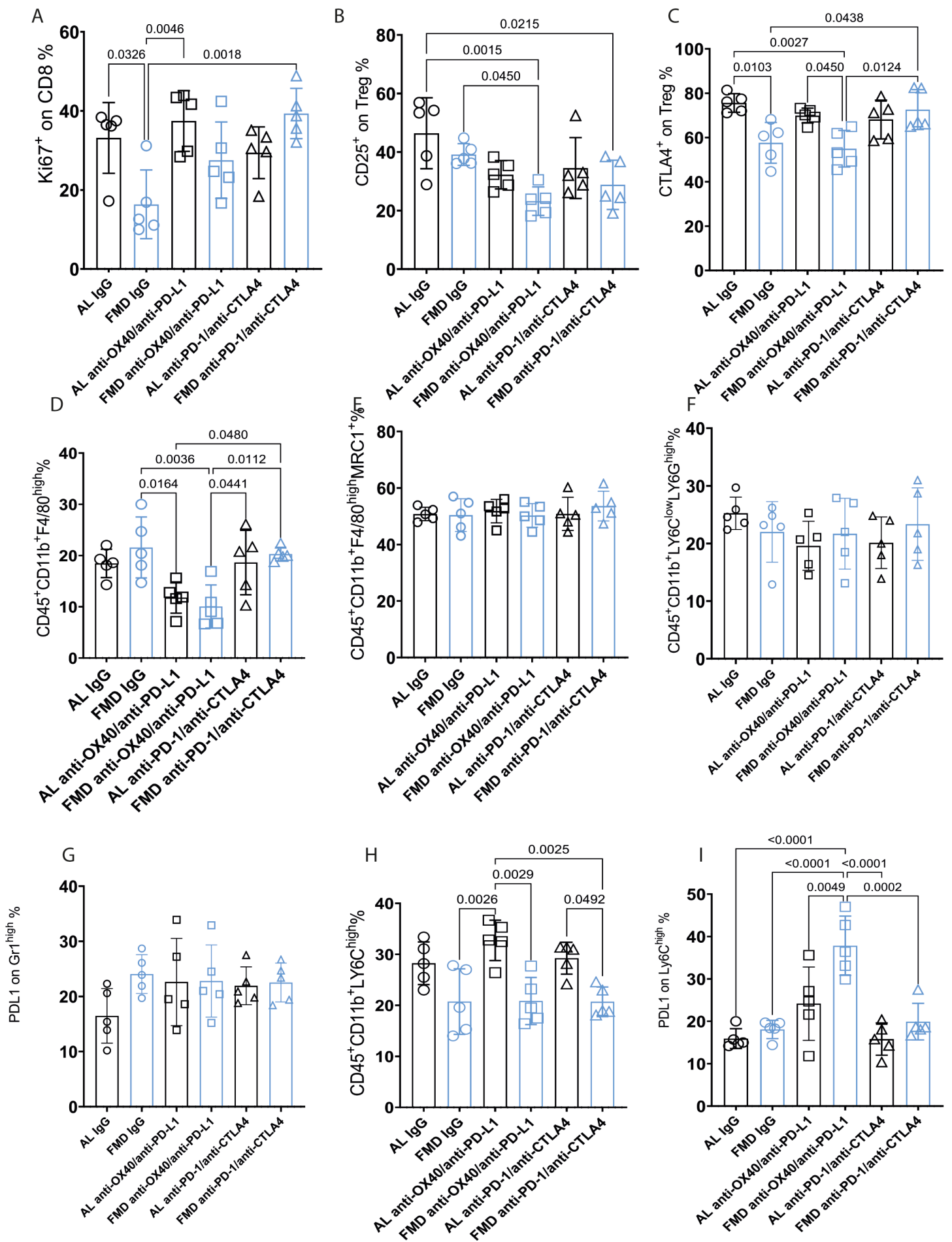
Supplementary Figure 2. IHC analysis of immuneinfiltrate in B16 melanoma tumor section. Representative images of tumor section stained with: A) CD8, C) CD4, D) B220, E) Myeloperoxidase. The images were obtained under 20 \times magnification. The scale bar was 100 μ m. B) Quantification of CD8 T cells infiltration in tumor sections. Statistical analysis was performed using one-way analysis of variance (ANOVA) (n = 3 tumors; 7 random fields/tumor). P values were determined by One-way ANOVA with Tukey's post analysis. Differences were considered significant when $P \leq 0.05$. All data are represented as mean \pm SEM.



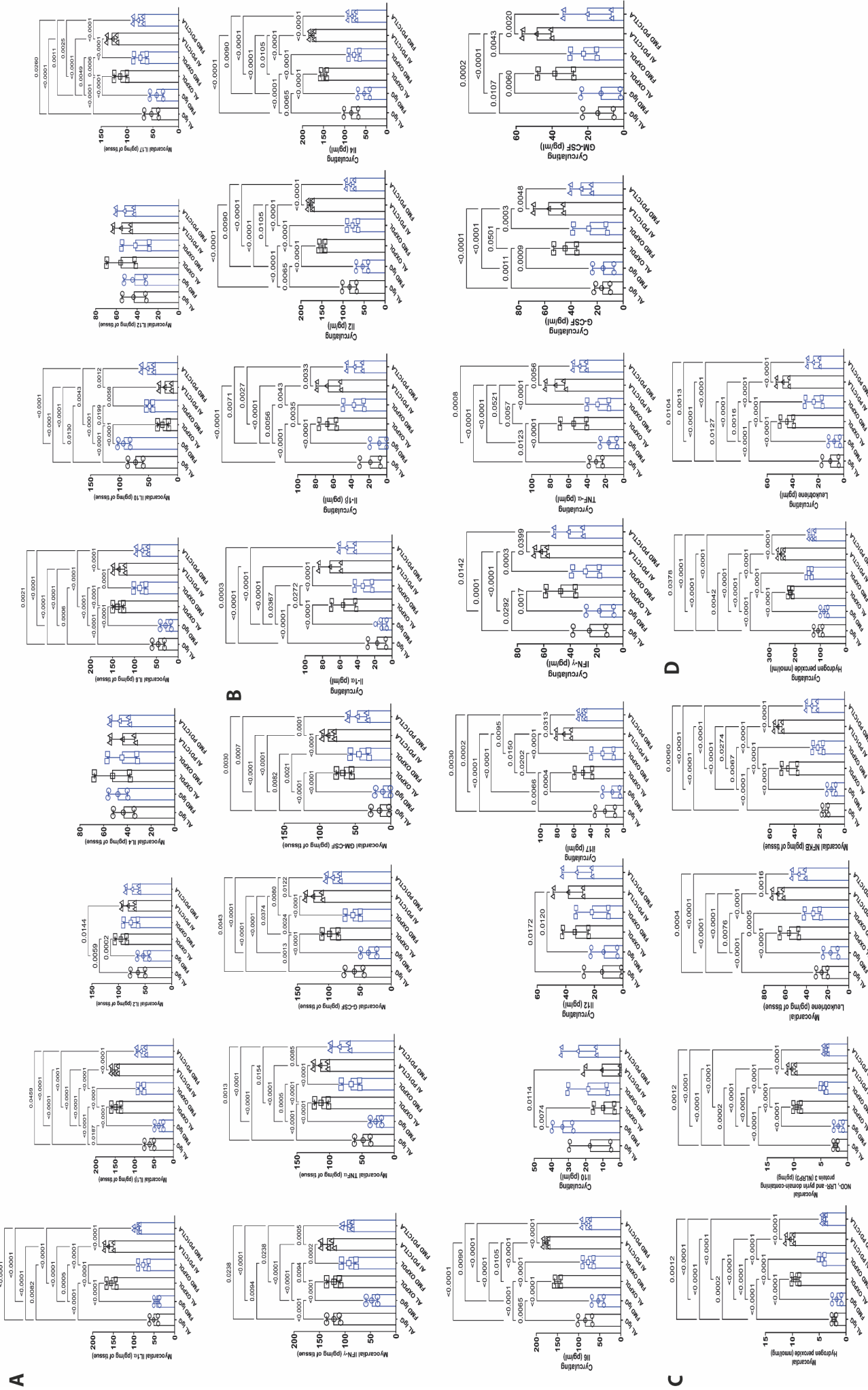
Suppl Figure 3. Analysis of B16 breast tumor infiltrating lymphocytes upon immunotherapy treatment in standard diet and FMD group. **(A)** CD45⁺CD3⁺CD4⁺ T cell (AL NT n=5; FMD NT n=7; AL anti-OX40/PD-L1 n=5; FMD anti-OX40/PD-L1 n=9); **(B)** CD3⁺CD4⁺CD25⁺ Treg cells (AL NT n=5; FMD NT n=6; AL anti-OX40/PD-L1 n=5; FMD anti-OX40/PD-L1 n=9); **(C)** mean fluorescence intensity (MFI) of CD127 in CD3⁺CD8⁺CD44⁺ effector T cells (AL NT n=5; FMD NT n=7; AL anti-OX40/PD-L1 n=5; FMD anti-OX40/PD-L1 n=9); **(D)** CD45⁺CD11b⁺F4/80^{high}CD11c⁺ (AL NT n=4; FMD NT n=7; AL anti-OX40/PD-L1 n=5; FMD anti-OX40/PD-L1 n=6), **(E)** CD45⁺CD11b⁺F4/80^{high}MRC1⁺ (AL NT n=5; FMD NT n=7; AL anti-OX40/PD-L1 n=5; FMD anti-OX40/PD-L1 n=7). Statistical analysis was performed using one-way analysis of variance (ANOVA). P values were determined by One-way ANOVA with Tukey's post analysis. Differences were considered significant when P ≤ 0.05. All data are represented as mean ± SEM.



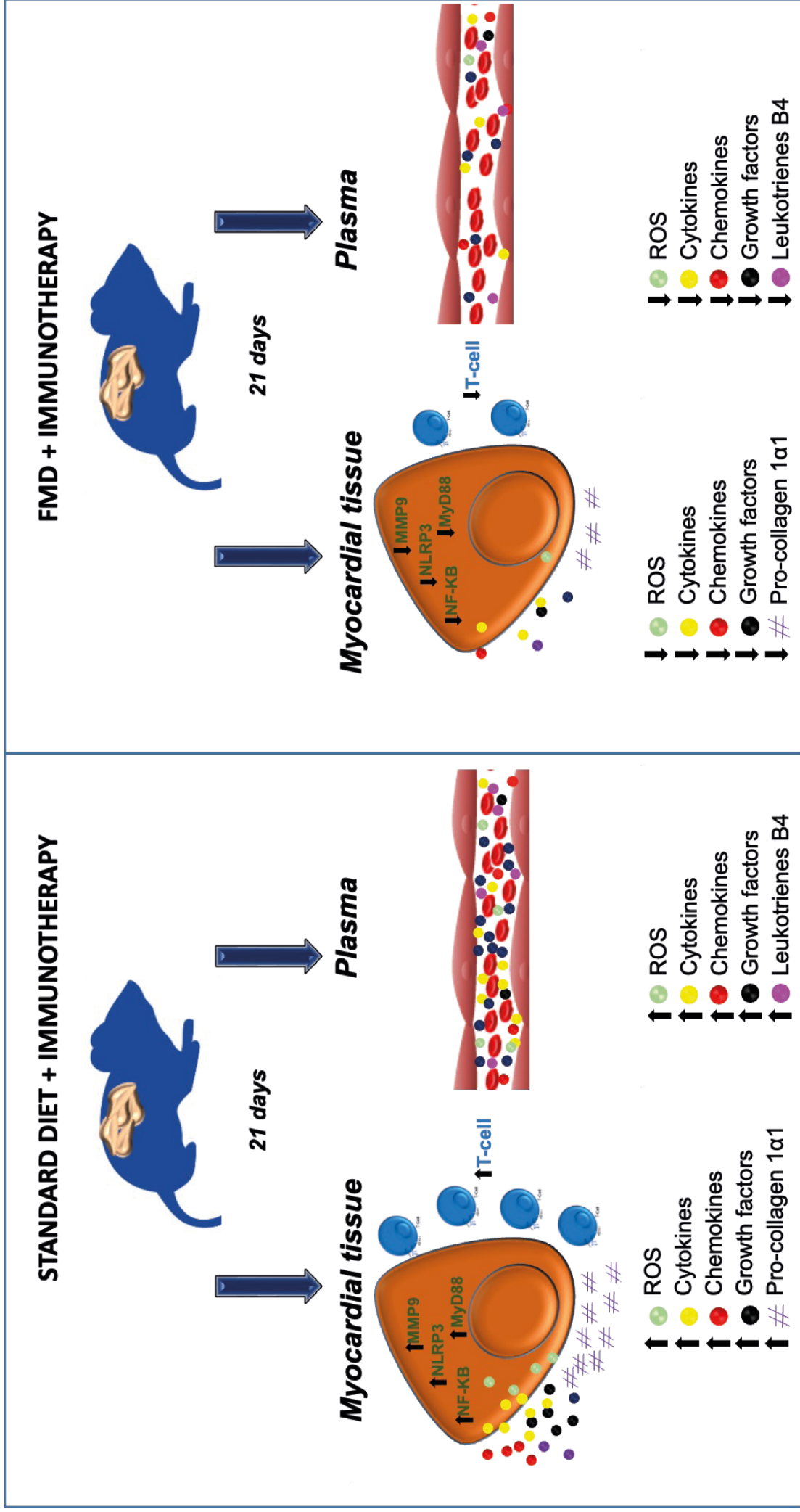
Supplementary Figure 4. IHC analysis of immuneinfiltrate in LLC1 lung tumor section. Representative images of tumor section stained with: A) CD8, C) CD4, D) B220, E) Myeloperoxidase (n=5). The images were obtained under 20 \times magnification. The scale bar was 100 μ m . B) Quantification of CD8 T cells infiltration in tumor sections (n = 3 tumors; 7 random fields/tumor). Statistical analysis was performed using one-way analysis of variance (ANOVA). P values were determined by One-way ANOVA with Tukey's post analysis. Differences were considered significant when $P \leq 0.05$. All data are represented as mean \pm SEM.



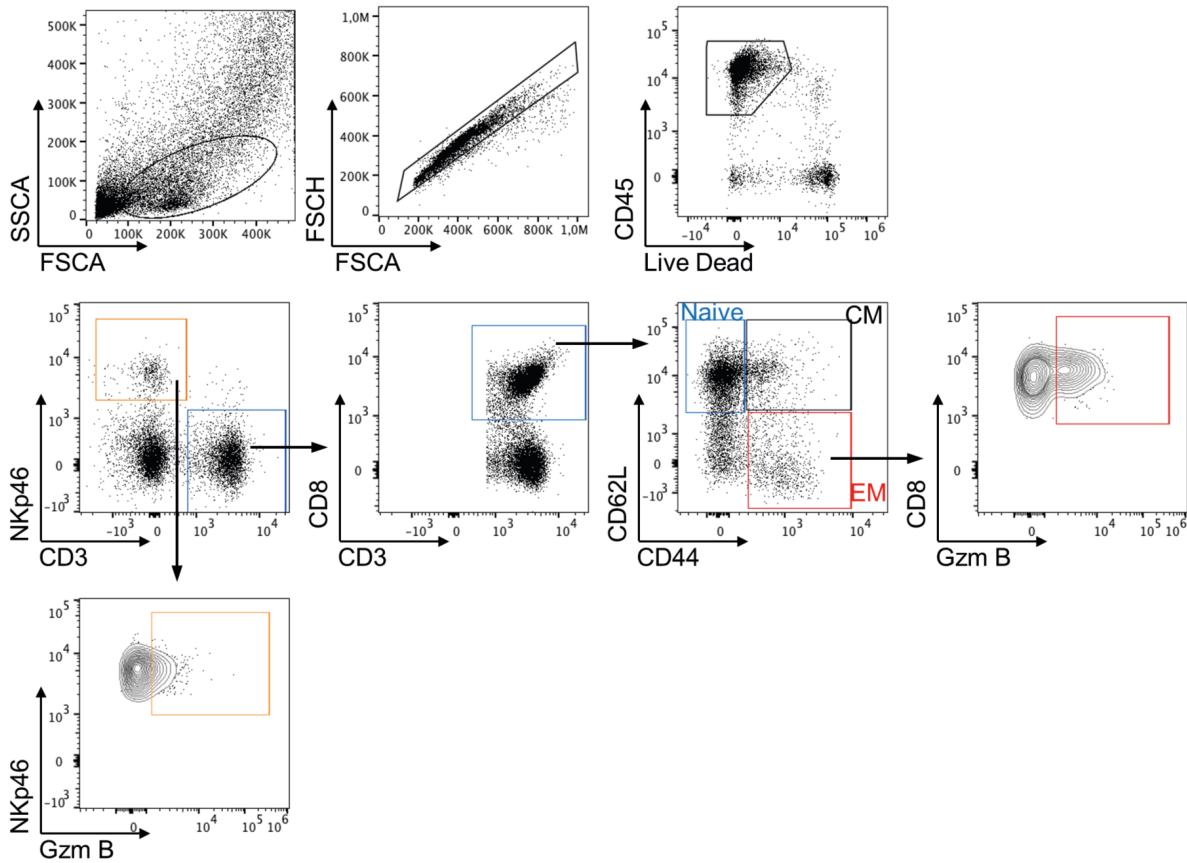
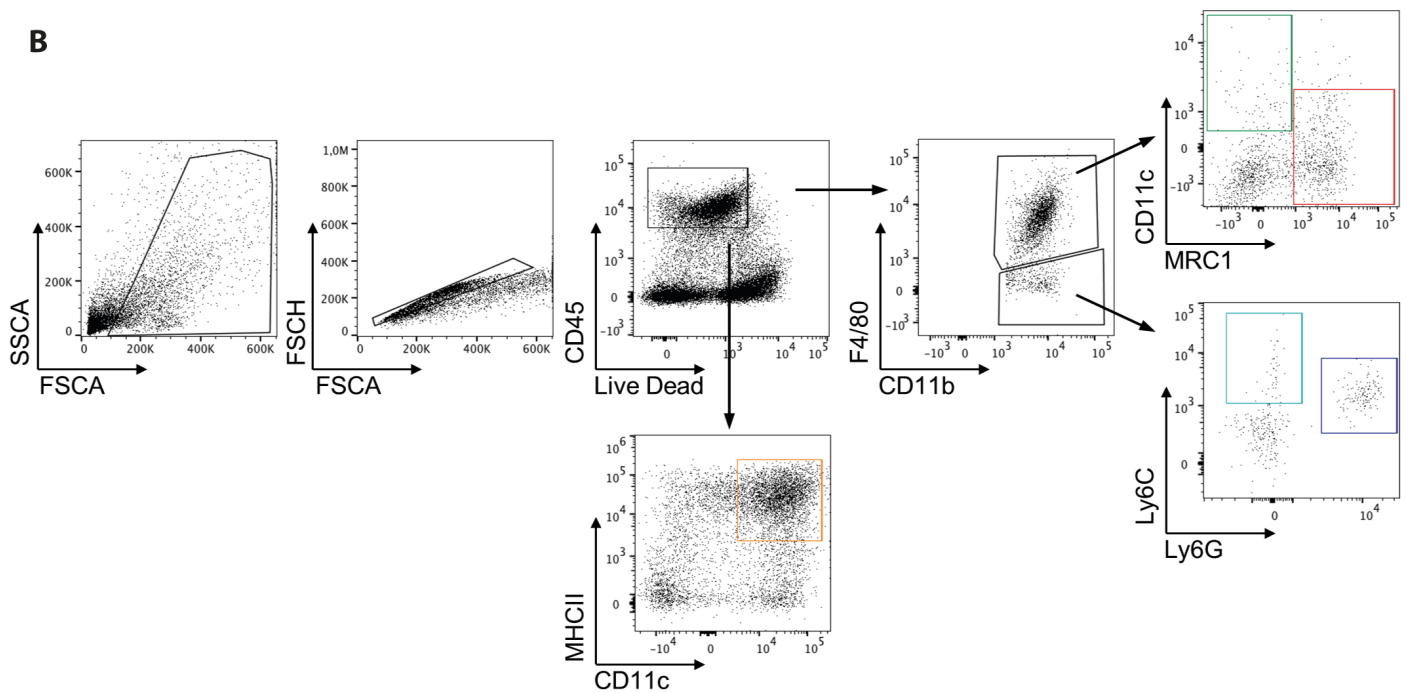
Supplementary Figure 5. Analysis of LLC1 lung tumor infiltrating lymphocytes upon immunotherapy treatment in standard diet and FMD group. A) Ki67⁺ on CD8⁺ T cells (n=5); B) CD25⁺ on Treg cells (n=5); C) CTLA-4⁺ on Treg cells; D) CD45⁺ CD11b⁺ F4/80^{high} macrophage (n=5); E) CD206⁺ on CD45⁺ CD11b⁺ F4/80^{high} macrophage (n=5); F) CD45⁺ CD11b⁺ GR1^{high} PMN-MDSC (n=5) G) PD-L1⁺ on GR1^{high} PMN-MDSC (n=5); H) CD45⁺ CD11b⁺ Ly6C^{high} M-MDSC (n=5); I) PD-L1⁺ on Ly6C^{high} M-MDSC (n=5). Statistical analysis was performed using one-way analysis of variance (ANOVA). P values were determined by One-way ANOVA with Tukey's post analysis. Differences were considered significant when P ≤ 0.05. All data are represented as mean ± SEM.



Supplementary Figure 6. FMD affects pro- and anti-inflammatory cytokine secretion upon immunotherapy treatment and attenuate Reactive Oxygen Species (ROS), NLRP-3 inflammasome, leukotrienes and NF-kB expression in heart and plasma. Cytokines level (pg/ml) in heart (**A**) and plasma (**B**). ROS, NLRP-3 inflammasome, leukotrienes and NF-kB expression in heart (**C**) and plasma (**D**). IL-1 α , 1 β , 2, 4, 6, 12, 17 α ; IFN- γ ; interferon γ ; TNF- α : tumor necrosis factor α ; GM-CSF: granulocyte colony stimulating factor; GM-CSF: granulocyte-macrophage colony stimulating factor (n=5). Statistical analysis was performed using one-way analysis of variance (ANOVA). P values were determined by One-way ANOVA with Tukey's post analysis. Differences were considered significant when P<0.05. All data are represented as mean \pm SEM. 'Source data are provided as a Source Data file



Supplementary Figure 7. Overall picture of the study demonstrating FMD decreases immune CD3⁺ and CD8⁺ lymphocytes and NLRP-3, Pro-collagen 1α1, MMP-9 and NF-KB expression in myocardial tissue of C57BL/6J melanoma bearing mice treated for 21 days with anti-OX40/anti PD-L1 and anti-PD-1/anti-CTLA-4 therapy. Systemic and myocardial chemokines, cytokines and growth factors(IL-1α, IL-1β, IL-6, IL17-α, G-CSF, and GM-CSF levels were also reduced in FMD group vs AI group.

A**B**

Supplementary Figure 8. Gating strategy for FACS analysis of immune infiltrate. **A)** Gating strategy for CD8 lymphocytes and NK cells (Figure 1 D-F and Figure 2 D-F). Among viable CD45⁺ cells, NK cells were gated as NKp46⁺ and T cells were gated as CD3⁺. Among T cells, CD8⁺ effector memory T cells were gated as CD44⁺ CD62L⁻. **B)** Gating strategy for myeloid cells (Figure 3 A-D). Among CD45⁺ cells, dendritic cells (DC) were gated as CD11c⁺ MHCII⁺ and macrophages were gated as F4/80⁺ CD11b⁺. PMN-MDSC and M-MDSC cells were identified among F4/80⁻ CD11b⁺ cells according to the expression of Ly6C and Ly6G.